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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/022,554A

Input Set : A:\PTO.VSK.txt
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#7
5-15-02
P.2.

3 <110> APPLICANT: ETTENSON, David S.
 4 BURGESS, Wilson H.
 5 DROHAN, William N.
 6 MACIAG, Thomas
 8 <120> TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO THROMBIN DEGRADATION RESISTANT
 9 FIBROBLAST GROWTH FACTOR-1
 11 <130> FILE REFERENCE: 054474-5005
 13 <140> CURRENT APPLICATION NUMBER: 10/022,554A
 14 <141> CURRENT FILING DATE: 2001-12-17
 16 <160> NUMBER OF SEQ ID NOS: 8
 18 <170> SOFTWARE: PatentIn version 3.1
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 638
 22 <212> TYPE: DNA
 23 <213> ORGANISM: Homo sapiens
 25 <400> SEQUENCE: 1
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 30 tcctctactg tagcaacggg ggccacttcc tgaggatcct tccggatggc acagtggatg 180
 32 ggacaaggga caggagcgac cagcacattc agctgcagct cagtgccggaa agcgtgggg 240
 34 aggtgtatat aaagagtacc gagactggcc agtacttggc catggacacc gacgggcttt 300
 36 tatacggctc acagacacca aatgaggaat gtttggctt ggaaaggctg gaggagaacc 360
 38 attacaacac ctatatatcc aagaagcatg cagagaagaa ttgggttgtt ggcctaaga 420
 40 agaatgggag ctgcaaaccgc ggtcctcgga ctcactatgg ccagaaagca atctgtttc 480
 42 tccccctgcc agtctcttct gattaaagag atctgttctg gtgttgacca ctccagagaa 540
 44 gtttcgaggg gtcctcacct ggttgacccc aaaaatgttc ctttgaccat tggctgcgct 600
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 50 <211> LENGTH: 154
 51 <212> TYPE: PRT
 52 <213> ORGANISM: Homo sapiens
 54 <400> SEQUENCE: 2
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 57 1 5 10 15
 60 Leu Pro Pro Gly Asn Tyr Lys Lys Pro Lys Leu Leu Tyr Cys Ser Asn
 61 20 25 .30
 64 Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly Thr Val Asp Gly Thr
 65 35 40 45
 68 Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln Leu Ser Ala Glu Ser
 69 50 55 60
 72 Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln Tyr Leu Ala
 73 65 70 75 80
 76 Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro Asn Glu Glu

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81	100	105	110
84	Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys Asn		
85	115	120	125
88	Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala Ile		
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106	tcctctactg tagcaacggg ggccacttcc tgaggatcct tccggatggc acagtggatg	180	
108	ggacaaggga caggagcgac cagcacattc agctgcagct cagtgcggaa agcgtgggg	240	
110	aggtgtataat aaagagtacc gagactggcc agtacttgcc catggacacc gacgggcttt	300	
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114	attacaacac ctatataatcc aagaagcatg cagagaagaa ttgggttgc ggccctaaga	420	
116	agaatgggag ctgcaaacgc ggtcctaaaa ctcactatgg ccagaaagca atcttggttc	480	
118	tccccctgccc agtctcttct gattaaagag atctgttctg gtgttgc acca ctccagagaa	540	
120	gtttcgaggg gtcctcacct gtttgcaccc aaaaatgttc ctttgcaccat tggctgcgct	600	
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137	20 25 30		
140	Gly Gly His Phe Leu Arg Ile Leu Pro Asp Gly Thr Val Asp Gly Thr		
141	35 40 45		
144	Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln Leu Ser Ala Glu Ser		
145	50 55 60		
148	Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln Tyr Leu Ala		
149	65 70 75 80		
152	Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro Asn Glu Glu		
153	85 90 95		
156	Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn Thr Tyr Ile		
157	100 105 110		
160	Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys Asn		
161	115 120 125		
164	Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala Ile		
165	130 135 140		
168	Leu Phe Leu Pro Leu Pro Val Ser Ser Asp		

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173 <211> LENGTH: 31
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175 <213> ORGANISM: Artificial Sequence
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178 <223> OTHER INFORMATION: PCR primer
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204 <223> OTHER INFORMATION: PCR primer
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212 <212> TYPE: PRT
213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: PCR primer
218 <400> SEQUENCE: 8
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RAW SEQUENCE LISTING ERROR SUMMARY
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VERIFICATION SUMMARY

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